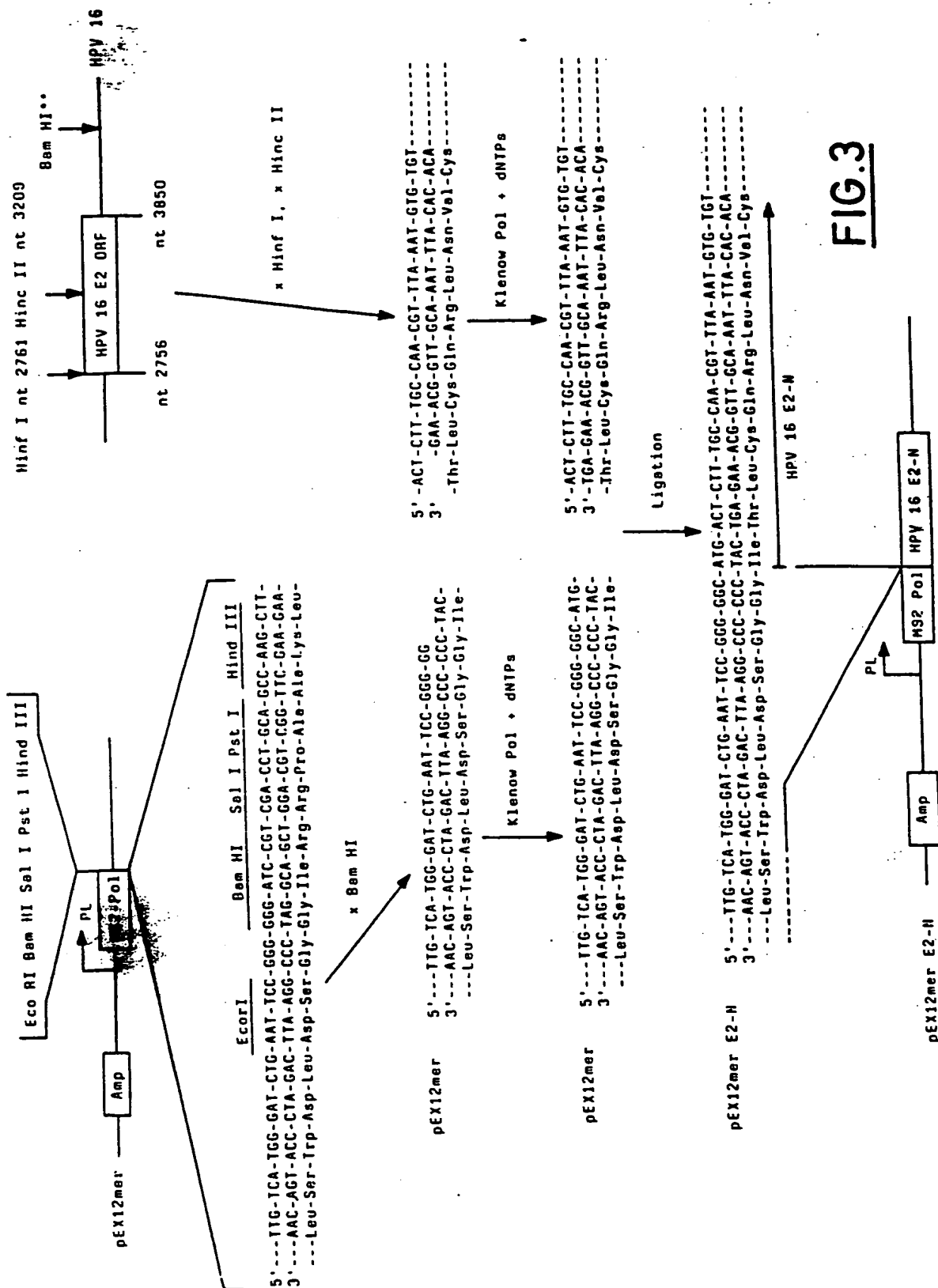


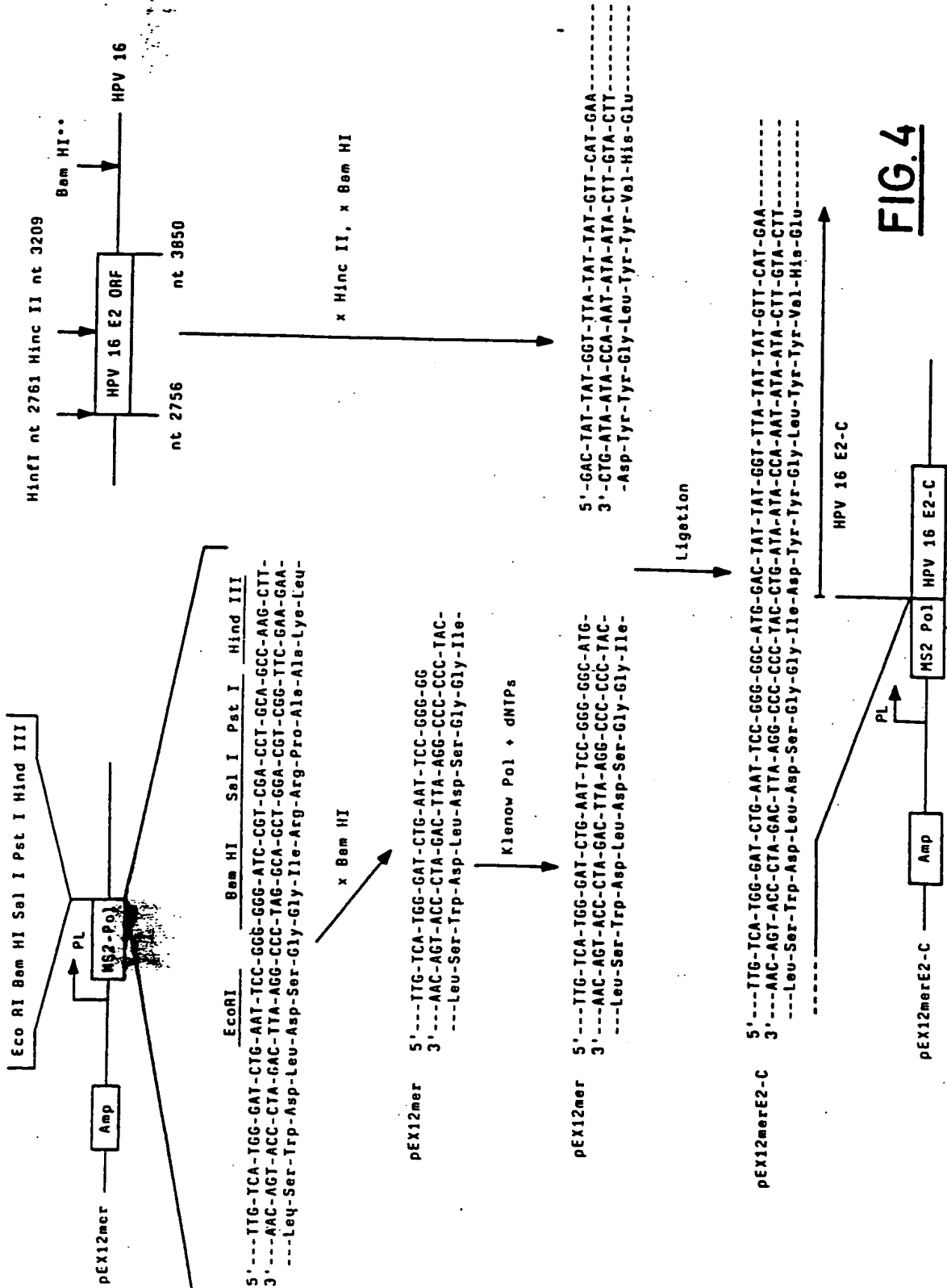


| Klon | Sequenz | Position | Isolate |
|------------------------|--|----------|---------|
| <u>Region E1-1090:</u> | | | |
| 1090 | NGWFYVEAVVEKKTGOAISODDENSDTGEDLVDFIVNDNYLT | As 16-60 | 1 |
| 1079 | NENSDTGEDLVDFIVND | AS 38-55 | 3 |
| 1084 | HADPAGTNGEETGCHGWFYVEAVVEKKTGOAISODDENSDTGEDLVDFIVNDNYLT | AS 1-60 | 3 |
| 1029 | EDLVDFIVNDNYLT | AS 46-60 | 5 |
| 1099 | EDLVDFIVNDNYLTOAETETAHALFTAQEAKOH | AS 46-79 | 5 |
| 1145 | NENSDTGEDLVDFIVNDNYLTOAETETAHALFTAQEAKOHRAOVVLRKRYLVVh | AS 38-91 | 1 |
| <u>Region E1-1059:</u> | | | |
| 1059 | stgsktkvfGSPISDTS | AS 92-99 | 1 |

FIG.2

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| Klon | Sequenz | Position | Isolate |
|------------------------|--|------------|---------|
| <u>Region E2-1066:</u> | | | |
| 1066 | DKILTHYENDS | AS 13-23 | 3 |
| 1025 | DKILTHYENDSTOLRDHI | AS 13-30 | 5 |
| <u>Region E2-1170:</u> | | | |
| 1170 | OLROHIOYKKN | AS 25-35 | 2 |
| <u>Region E2-1074:</u> | | | |
| 1074 | AIYYKAREMGFKHHNHVVPTLA | AS 41-63 | 4 |
| 1060 | AIYYKAREMGFKHHNHVVPTLAVSKNKAL | AS 41-70 | 2 |
| 1061 | YYKAREMGFKHHNHVVPTLAVSKN | AS 43-67 | 1 |
| 1063 | INHVVPTLAVSKNKALQAI | AS 55-73 | 1 |
| 1057 | INHVVPTLAVSKNKAL | AS 55-70 | 4 |
| 1018 | TLAVSKHKALQALTELQTLTETIYNSQYSNEKNTLODV | AS 61-76 | 2 |
| <u>Region E2-1112:</u> | | | |
| 1112 | ULTLETIYHVSQYSHEKWTLODVSLE | AS 76-100 | 1 |
| 1156 | TLETIYNSQYSNEK | AS 78-91 | 1 |
| <u>Region E2-1158:</u> | | | |
| 1158 | TSVFSSNEVSSPEII | AS 197-211 | 1 |
| 1121 | VFSSNEVSSPEIIROHLANHPAATHKAVALGTEET | AS 199-234 | 2 |
| <u>Region E2-1102:</u> | | | |
| 1102 | EIIROHLANHPAATHKAVALGTEETOTTIORPRSEP | AS 209-245 | 1 |
| 1091 | TEETOTTIORPRSEPOTGN | AS 231-249 | 2 |

FIG.5